


# Comparison of mutation severity, gene intolerance, and mutation distribution

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 An abbreviated version of this protocol was published in Science Advances in Jan 2022


De novo mutations identified by whole-genome sequencing implicate chromatin modifications in obsessive-compulsive disorder

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## Detailed protocol

Please see the attached file for the script.

## Related files

 protocol\_distribution.txt



**How to cite:** (Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Lin, G. , Song, W. and Wang, Z. (2022). Comparison of mutation severity, gene intolerance, and mutation distribution. Bio-protocol Preprint. [bio-protocol.org/prep1704](https://bio-protocol.org/prep1704).
2. Lin, G. N., Song, W., Wang, W., Wang, P., Yu, H., Cai, W., Jiang, X., Huang, W., Qian, W., Chen, Y., Chen, M., Yu, S., Xu, T., Jiao, Y., Liu, Q., Zhang, C., Yi, Z., Fan, Q., Chen, J. and Wang, Z.(2022). De novo mutations identified by whole-genome sequencing implicate chromatin modifications in obsessive-compulsive disorder. Science Advances 8(2). DOI: [10.1126/sciadv.abi6180](https://doi.org/10.1126/sciadv.abi6180)

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